

DATE: 01/29/2003

TIME: 15:14:26

OIPE

# RAW SEQUENCE LISTING

FATENT APPLICATION: US/09/830,913A

Imput Set : A:\pb0005usn\_seqlist.txt
Output Set: N:\CRF4\01292003\I830913A.raw

```
4 <110 * APPLICANT: INCYTE PHARMACEUTICALS, INC.
              WALKER, Michael G.
              VOLEMUTH, Wayne
              KLINGLER, Too M.
      9 (100 - TITLE OF INVENTION: CORTICOSTEROID SYNTHESIS-ASSOCIATED GENES
     11 - 138 - File Reference: PB-0005 PCT
C--> 13 <140> CURRENT APPLICATION NUMBER: US/09/830,913A
C--> 14 <141> CURRENT FILING DATE: 2003-01-29
     1. Hittle Paice Application NUMBER: 60/155,269
     17 :1-1 - PELIS FILING DATE: 1998-11-06
     1 4 -: 160 - NUMBER OF SEQ ID NOS: 9
        :170 GOFIWARE: PERL Brogram
             SEQ ID NO: 1
     .4 :211 · DENSTH: 3736
              TYPE: DNA
     16 -2013 - 08.GANISM: Homo sapiens
     LA - ... G FEATURE:
     . * - DL1 - NAME - KEY: misc_feature
     -4 - 228 - DIHER INFORMATION: Indyte ID No.: 0649730B1
     PI HANG SEQUENCE: 1
         -totiltopagt tugagaagag gagtiggtgat aaagoobact gacottoaca battotaaaa -60\,
         witggttggo aktgccagta taccaacaat catgatcago tgaaagaaac agatatttta 120
         ustigopaga abaadaaatg atgapgoaad tabagtbaga tetatagoda godatotato 180
     50^\circ aponotagas gottocagag acagugaaao typaagatyo totoaacagg attatytoto 240^\circ
         atggagadda gtaagaaaat batttatotg aaggtgaaat gbagagttgg ataagaaata 300
     38 -pathyptiggs thichasast getydotitod typototapt obsobiodat bootggapti 3.00
     50 -throughoutly gradiagrage bluappadet toacceptigt geaccaceca agasagages 4.05
     40 -alabettiged throsactity galaxtyotgy ggt bootggt gtggtaagaa actosscate 4\%
        -arabgggtat ghagaaggat goobtootgg gatttgbagg tabataaaaa atgtatggba 540
     40^\circ touthtoott ghasattott phagtitopa agtgagaagg ggagbaggtg titactgatg 6\%
         \eta_{	ext{AdA}}agytat gitgotatgt tyatqtgtaa qtqaaatcay ttqtgtgcaa tagabagggg 600
     4.7
         -butiaticatic dyadoaticad ocayitticta sasoccadag godaticagos gotagaggitg 7.\,\%
     .; .;
         ymt_{4}gottt_{6}godagadatg gaddotaaat baadagadaa t_{9}goatt_{9}to gaagagdaad _{7} ^{10}
         styttaatpa atsatyttaa aaatsaaggt ttygottoay titaaatsas ttyaggtaty 841
         algorithator by this body agata about adjuty atomic top calcate and acceptage 9\%
     .; 7
         questatear a Habbatoad tagitization by:tagitize by tagitizating interesting 9n \phi
     . . . . .
         twaagobatq babbabagab ttotgggbag agotgagaga baatggtoot gabataataa 1 20
         gyatottiga traaccocca taaggoatgi yigigiatac aaatatacti otottiggoi 1930
         withogapata gaadotbago tifttaadbaa jiffgaaatab atbagatotif baababagaa 1140\,
         Abgribbiged transitione abbatgeota gyantosepp battitateda gytottibity 1.00\,
         gatingstia arcaminage estataatea etigetamas asiggetie atemeesagg 1.60
     . . . 3
         gataaaaaca gigatsattg tittiggadot ootgoatday istattoaaa attatotitto 1520
         tototagott tunacaaato otaaaattoo tytoocaayo cacccaaatt oteayatott 1380
```



FATENT APPLICATION: US/09/830,913A TIME: 18:14:0

PATE:

Input Set : A:\pb0005usn\_seqlist.txt
Cutput Set: N:\CRF4\01292003\I830913A.raw

```
ttotggaada aggoagaata taaaataaat ataqatttag tggottgggo tatggtotoo 1440
      aaagatoott baaaaataba toaagobago ttoattoabt babittabtt agaabagaga 1800
     tataayggoo tgggatgoat tfattttäto aataobaatt tttgtggood tggodgacat 1860
59 tgotaatoaa toasagoast atttootatt aagoosastg atttottoas aatoottoto 1620
80 -aaattacaat teeasagago ogobactosa bagteagstg aacceasbag teagatgaga 1680
\sqrt{1-} gauatqsaco etaettyeta tetetatett agasageaaa sacaaacagg agitteeagg 174\%
\kappa Z -qagaatiggga aayibaggga goataaaaaga tabagtibagg gaaaataga totaggoaga 1800
63 qtgcottagt baqqqabbac gggogdtgaa totqbaqtqc baabacbaaa otgabacato 1860
64^{\circ} throaggigta peticpaabbe tagbottoto beacagoige blacaacaga gibbobbago 1920
     ortotoagag agotaaaabo agaaatttob agactoatga aagcaacbob ocagostoto 1400
\mathbf{r}_1 \cdot z
      \phi meansoring organizated that the transfer and according to the state of the second of the second second of the second of the
100
      thataagbag gggbbagaat abobbagbba bobgbagtga babbgbtgga bobbtgaaaa 1000
60
     -mrattopata qqaqaatqqq ttooobaqqo toabaqtqta qaqabattga qoobatbaba 1101
\mathcal{C}^{+} autiquitinga ofgotopolog totalaaldag todaloodago obatggoladi googogtgat \mathbb{A}[\mathbb{A}]
Which topqoqqoo attoaqaagt toaagoogag atgotgacgt tgotgagbaa ogagatggtg \omega \ell
Ti arcatoagtg casatgoado attoagoada toagtoatat godoagtgoa gttadaagat ..:40
      quequitogg dasaqoatit itgatggaata gggaactgia satgtatgat gattitigasa 249\%
      Aggotbagoa ggattigtto tiaaadbgab tbagtgtgto atbobbggtt atttagaatt 1400
     -anagttaaga aggagaaact totataagad tgtatgaaca aggtgatato ttoatagtgg 1^{
m su}C
     gotattadag gdaggaaaat gttttaadtg gtttadaaaa tooatdaata ottgtgtdat 1950
TW -topotytaaa agybagyaga batytyatta tyatbayyaa abtybabaaa attattyttti.940
     -thagecodeg typicategic efficies gradet greateritti fitattaaage caaatetgig arepsilon^{\prime\prime}
78 engbatatab togbattoba tgtgttagat ggaagdattt optatobagt gtgaataaaa . 😘
Fig. Agaacagtig tagisaasta tiataaagoo gatgatatti batggoaggi tattotabba DF20
80 -agotytyctt gttgyttett oodatgadty tastysttt ataaatytas aaatayttas i88\%
     -typapatgang agandottgt tignanagna tiaataagaa cottgataag aancatatto 1.94\%
      tyttyadago dagotdabag titottydot gaagottygt ybabbotboa ytgagadaba 2000
      arabototot titaboaaag tigagaacag agoiggigga tiaattaata gibtiogata I(\theta)
1.5
64 thtggodatg ggtaadotoa ttgtaadtat batbagaatg ggdagagatg atbttgaagt 1000
     Ambubataba bisaaagibba aababtaigi bagaiggggg taaaatobat taaagaabag 1960
     grasiastas tratasgatg atasgosast gtitosgoco astgicasoc cagitasasa NAC
AV lasaauttaat gotgigtaaa atggttgaat tagtttgcaa actatataaa gacatatgca PPK
      ighaaaaagto tyttaatgoa catootytyy yaatyyayty tootaaccaa ttycotttto 1960
44 tightatotg agototocta tattatoata otoagataac caaattaaaa gaattagaat 1420
90 andantitta atabaettaa eattaaaete tietaaetit estetiotg igataaitea 3490
AI -quadrataget abggatoses autgeototic agreeatiget abadaaaats agetatodot ^{10}\,\mathrm{M}_\odot
      ataboatgot ataggagadt gggdaaaaoo tgtabaatga baadootgga agttgotttt 3:000
4
93 titasaassa tastasatti ottaastosa otottittio iggitytoig titgitatas Prof-
(4) automasogt attosagtod tosatatoot gatostaata oostgotata ggagaotggg 37.00
     -baadabbtgt adaatgadaa obotggaagt tgbttststa aaaaaaataa taaatttott 1780
                                                                                                                       3736
196 - Agatomadaa agaada
93 <310 - SEQ ID NO: 2
100 -0011 - DENGTH: 1187
111 - L1L - TYPE: DNA
1/2 -11: ORGANISM: Homo sapiens
1.4 -1.20 - FEATURE:
1.5 · 1.21 · NAME/KEY: misd feature
196 - 121 - OTHER INFORMATION: Incyte ID No.: 965781CB1
108 <450 / SEQUENCE: 2
```

### RAW SEQUENCE LISTING

FATENT AFFILICATION: US/09/830,913A

Imput Set : A:\pb0005usn seqlist.txt Output Set: N:\CRF4\01292003\I830913A.raw

CATE:

.1-24 .1...

71%E: 18:14:5

```
-atalappatt astalapyja titygoobto gagydaagaa atloggoasy agyalbagao -6.
          catcagaagg attigiataa agagigasis isotaigaag giaaaggesa suosistiva 120
          gitocagiga orgagatada ittitodaat ootgggggda aatacagaba dagbaagtto 180
         offictions, figgaaatti ggoagetgee ticadcagtg agcacaaage bacatticaa 24\%
     113 aggaaactga caaattates ecagetgoba gaagaagaaa tootoactga acggottoot.
         gitteetyty yileattate iqattyyety cagyyatyaa agittitaay tibalaggac 360
         tgátgáttet esteácetet gegítttésag edgyttésagg ácaaagtoga átgástytge 420
     116 tgtgotorat agactygtto atggtoacag tgcacccctt catgetasac aacgatgtgt 480
     11...
         gtgtacantt teatgaacta bacttggged tgggttgeed bebaaacbat gtteagodae 540
         .updoptarca gitcapotac ogigtiacig aaigtggdat bagggddaaa goigicicic 600
     115
          gagainatiggt tarbotabago abbgagatab abtabtotto taagggbabg bbatbtaagt 600
     119
     1.2
          tigityatood agigtoatgi gotgooodd aaaagtoodd atgydtoadd aagoodtgol \mathcal{T}(0)
         poatgagagt agodagbaag agbagggoda bagbobagaa igatgagasa tigotadjagg 781
     101
          tyttoagott gibabaytoo aytbaalaggo bbaabtybga ttytobabbt tytytottba 841
         ytgaagaaga goatacccag gtoocttgto accaagcagg ggotcaggag gotcaaccto 9.1
     1. :
         tigoagopato toaotttott gatatttotg aggattiggto tottoacaca gatgatatga 900
     1. -
     128 htgqqtodat gtgatootba ggtttggggt otdotgaaga tgotatttot agaattagta 1920
         , hatigitatas akaiqitoiga baaatkagig ototigigas poidaigiga goadittiga 1.80\,
     1. .
          gaaagagaaa ootatagbaa ottoatgaat taagoottet totatatett tatattoatg 1140
     1.
         tgtwaabaaa awataasata saattotgat ogostaasaa aasasaa
                                                                               11.67
     1. -
     1:1 :::14 - SEQ ID NO: 3
     1). RULL - LENGTH: 1395
     133 - Mili - TYPE: DNA
     134 -0013 - ORGANISM: Homo sapiens
     137 -0.020 - FEATURE:
W--> 137 <221> NAME/KEY: misc feature
     136 HA23 - OTHER INFORMATION: Incyte ID No.: 1419725081
     140 -:400 - SEQUENCE: 3
```

140 otokasggos akossasaggs katgoboggo tooboatggo tyttgobayo soottostab-6%-usgrigotoss opotosgaso potostitata toposagigita agastoposas titagogotos  $1,\,0$ 14. -orgatigaty digggagtag gggtataggg tiggaggaatt aagatgatta gagaatgggg  $1\pi \delta$ 14: 144 -scalagoggae aggstiggtit gggaggetga ggtototatg gtiggtiggatg gaaaaggaga 74%Agosagggao tagtgagada dogggtotgo obcaaagott tgtgttgtto totopatggg 300 1::: 140 -aacagggtaa cagcagagga aggatogaaa goocaggcag agaacttaga aatgcctgct 560 14/ onalatatat ahantgotga gtopagaaag atdagggott dagaggtotd byottotgod 450 totogggatg goggggdagg gogattotog tgagtgggag agtotgtgot gdaggtadad 480 14. 149 tggabactgo bbactactto ggttoagogt gggobagado tatgggbagg tgabtggtoa 540 Tatwottogs ggoodlooty gootagooty goodcotyto dadegeadad ttotgooted 600 1:0 1:: -pattoggoot obsagatoto otgaggttoo pagggagagt otapotgtoa ggogtgggoa 660 15. Agahaggoto agotocagoa tgatocotgg gtacacaggt ttgtacgcag gtatgcacag 720 Tripopostoc caggageaty ryttocaaac actaacgagt ettecttyte cetycotyce (%) 1:.: -bagusettyt abbooggyba bagttbatot styppaagaa otypagopag gtotyggybby 64%154 aggetotgag tgaptttabt babttgbatg aaaagbaagg gagtgaagag btabbaaagg  $9\omega$ 155 1: • aga seaagan aaraaandab acaqaqaaad accaggtgoo agagooggan niggoagotgy 900 13.7 Aggagoogad actggaggtg gtggaadaag ottotoocta otocatggat gadagggade  $1000\,$ 185 otoggaagit ottoatgica ggdactgdag gaattigggd agaagdacto accaggdagt  $1.60\,$ goodaggado odasabatot odoodaatti oodagaadat adootoagaa ootgggtott 1140 15. thapphaaph abgggggta ogtgobaggg bataagthop aghtbggbba babathtggb  $13.90\,$ 100 catetoacco atgatgetet gggeeteage aeetteeaga ageagetett ggettaggee 1200 161

IAIE: "Isla I

```
TIME: 15:74:2
                      FATENT APPLICATION: US/09/830,913A
                      lnput Set : A:\pb0005usn seqlist.txt
                      Output Set: N:\CRF4\01292003\I830913A.raw
         - actidascáto máditecent pecititical equatocomo ecatrotitit gyadydyaga 1310
         - gaggtgggtg ggagggtggg ggaababaaa gagaaaatgg tittggagget gagpabbitt 1880
     164 - titattaata ggiat
     167 K2185 SEQ ID NO: 4
               LENGTH: 36
     168 <211
     169 K211 - TYPE: DNA
        ~213 ORGANISM: Home sapiens
     173 < 22) + FEATURE:
W--> 173 <221> NAME/KEY: misc feature
     174 KA23 - OTHER INFORMATION: Incyte ID No.: 23645820B1
     176 -: 40 - SEQUENCE: 4
     177
          stiltetrag basagygdag daetgetege tittgetyggg dagedetede aggaagated (6)
     17.
          -bawaagahat gototgaggt gigigaabab igigaatgib aiggabaaag bibagogigg 1/\phi
     173
         -gaouscatgy gusticaggs obtgaaagsa goagatotba botgbattgy aggatgagas 1:\mathbb{N}
     180 genecatyaa gwagaaagaa gabgobabag abagbagbagbab aytabbabot tobagagata 240
     181 grateactyt gadactbagt tiggaatyca atygogogat bitggotcae tyaaaootoo 3.0
     18. Hettoppagg tilbaaabgat totootgoot bagoottoog agtggotggg abtabaggaa 860
     18: Annagotyto aggostoppa gatagtatba gaaagotyaa gatttopaga togotypato 410
     184 Intercaphas appropriates operquented actacotact gootyttyag casotoctac 440
         -alwasactoot sacttratty gitabagaga cagattitag abitgiotoc batatootyg ^{1.4\%}
     18
                                                                                 5.67
         studatoadd agdaataaag dottadt
     18 \, \mathrm{m}
     189 KD17 - SEQ ID NO: 5
     190 -: 111 - LENGTH: 920
     191 H. L. TYPE: DNA
     191 HL13 - ORGANISM: Homo sapiens
     194 H. ST. FEATURE:
W--> 195 <221> NAME/KEY: misc feature
     196 HIGG: OTHER IMPORMATION: Incyte ID No.: 2737624081
     19: -0400 - SEQUENCE: 5
     199 -gog-salabot ghagtegtag obabttggga ggbtgaggba ggggaatbab ttgaabbbbg \sim 60
     \Omega(x) -ray reggagg thirtainiggs cogagathyt gecaphges becageging gegacagagt \Omega(x)
        - daduoticoat oficaaaaaaa aasaaaaaaa aatotatgot agtagattad aacttcacac 190
        - tayaggaytt otygabasay ottotaatta gtosasotsa sitasygoto sitassagga 14\%
         - waqqaactab tiggaaatta tigbaattoaa taatttagac totgttacca ggatctttca 🥬
     \pm 0.04 taasaaattta aittopataa toataabota aatgagttot taaaagaatto tataagbaat 3\%
     201
        – agetgattaa teeggeeergg aagatgaaga tbabaactgt bbabbbaccb aabbaaaagg 420
     \pm 600 aalgegbagty chaalatatga gaggataaad aatattaytt aabatttoty toatttatga 480
     N(\mathbb{N}) typomaattay tagtaayata attomadago tytomamitt ytttyygyst yymmattitt \in M(\mathbb{N})
         -tighttasac akgotsasag totagtatto tgggagaagt ggotggaaga aggggtaata \epsilon \in \mathbb{C}
     _ () ~
     \pm 0.8 tayigaaago awattoocii toocaaggagt caaqagaatt taigigaagi ggotoocgoo k\pm 0.0
     110 tgtaatotoa gosotttggg aggodaaggo aggoaatoad otgagggdag gagaddagdd 710
     111 tgüctaacat ggagaaacob catototaot aaaaatacaa aatgagotgg gogtggttggt 280
         - qoangoctgt awtochagot acttgggagg otgaggbagg agaatogoot gaabooggga 840
     ILE ogoggagutt gasgtuages aagategage caetgtaete cageetgggg gacagatega 🔫 0
                                                                                 920
         - agotgtotta aasaasaaaa
     :17 -:21% SEQ ID NO: 6
     Lib -: Jill- LENGTH: 1564
     21" W. 11 TYPE: DNA
```

RAW SEQUENCE LISTING

220 KL139 ORGANISM: Homo sapiens

### RAW SEQUENCE LISTING

FATENT AFELD ATTIME US/09/830,913A

Input Set : A:\pb0005usn\_seqlist.txt
Sutput Set: N:\CRF4\01292003\I830913A.raw

:ATE:

T1ME: 17:14:.

```
AZZ WILL FEATURE:
          NAME: KEY: miso : 4: ture
224 Kalas Pother information: Incycl ID No.: 2860066081
226 *400 - SEQUENCE: 6
    ammitti aa aasti hotto tiisaa git toalgatta. Abatigilig baaagtaaa.
    - gravosagott samitinotym argottämit tyrimagasi tatatinimat samasamity 12
    ilm pubagang war nabagon ipigobranos graatgagna tabah ging garibpbagai libi
   tytuttiong gognorgoan apgeaagnee craabtegen gaggagaget gitteagenat 240
    ggorgotyte acabattaes tytatetity emagnitage tygatyetea tteagtetyt 300
    gaarttotyg taogtystyg tyatgaatga tyagoasaca yagagysgat atotystytt 360
    thinditidig agingyggad taddagditt tgiggigatt diddicatag tiattitgaa 420
.334 aggmatotat batbagagda tgtbadagat otatggabto attdatggtg acctgtgttt 480
    tathocasad geotargoty effigitioad typagetott getectitya ogigeologi 540
    ggtagtgtto gtggtyttow tocatocota ppaggtgaag ppacagtgga aagcatatga 600
.13€
337
    tgargtette agaggaagga caaatgetge agaaatteea etgattitat atetettige 660
   - totuatttoo gtgacatggo titgggggagg actacacatg gootacagac acttotggat 720
.339 gutugitoto telgipatel loaabagist gbagggasti talgittica iggittatit 780
     -satificadad aadbaaatgi giigoddiai gaaggobagi tabacigigg aaaigaatgg 84\%
..41 goalootgga bodagbabag botttttbab godogggagt ggaatgbotb otgotggagg 900
    - igawatcago aagtobabbo agaatotoat oggtgotatg gaggaggtgo cabotgabtg 960
    . Igadagagoa toottobaab agggbagtba ggobagboot gatttaaago baagtobaba 1820
- Haatigaagoo abgotboogo boobotggagg atatggobag gygobabtga taybogatga 1\%0
. .
    - qqaqtboobaq qaqtttqatq atttaatatt tqcattaaaa abtqqtqbtq qtbtcaqtqt 1140
    -bagigataat gaatotggto aaggoagoba ggagggggo abbttgabtg abtoobagat 1270
... d 6
    -agradagata aggaggatab spatogooga baatbabatg tagbabbta otaacbatto 12\,60
347
    yabryagbab abitibatat tigiatbago tittyigota aaabibibta agiabatbba 1920-
    optytytaat aggaapotyt gaattytabt ygatyattaa tabaaaoyty attyttytat 1340
    ntgragtata aattactgat tgtatgtgad otgaaaatto actgotataa gaaaggtgga 1440
    jublightigt ambagstaat aggatgttba tättbbaagg atattagstig titttttaat 1810
    pateotatat gyotaabatt giitaatgaa agiaataato aataaagbaa tagaatotaa 1600
    Hāda
1366 <310 SEQ ID NO: 7
.57 <211. LENGTH: 752
1188 <31,2 × TYPE: DNA
HER KA13 + ORGANISM: Homo sapiens
281 K223 FEATURE:
202 K221' NAME/KEY: mist feature
263 K223 - OTHER INFORMATION: Incyte ID No.: 29615630B1
365 <400 SEQUENCE: 7
\mathbb{R}[66] gatqgqagqg ggtagqgagq atqaaaaaat baccattacc abbtqaaqbt titboattot 60
NOT light-thaata goattyatga titgdaddat tiggagatag agototygot tatttattat 13
\pm 60 againsaggag stigggestag tiggtigestad scalesatiged agestactotig aggistigagg 1\%
北京 tgggagāatt gotggagggē aattogaaac bagtotggge aagatagtga gabotbatot 241
    caaaaaaaaa aaaagaagga aaaaacaaag gagttgatga aaccaaatta tgacaaacta 311
270
   aattaaaaaa ttaaaaaata goattagtgg gaacactgtg gtatttootg aggtattaaa 361
272
   attittacaac tagtaaaggi attiattata attiattogg attyttott agittattii 42
273 tgacatggat aaaattattg tttatctagg agootcagag ttgagagagg atagttgtag 45
374
    agaagtatti otaaaatyga attatoatoo actaccagot aagagaaaat agitactig( 54)
275 - ttaaagatac ottaaagsaa datatattgt agtttoaaga gaasaggetg acagtcaaga 600
```

# VERIFICATION SUMMARY

HATENT ABBILINATI N: US/09/830,913A

Input Set : A:\pb0005usn\_seqlist.txt

Cutput Set: N:\CRF4\01292003\1830913A.raw

L:18 M:070 f: Current Application Number ditters, Replaced Current Application Number D:14 M:071 f: Current Filing Late ditters, Replaced Current Filing Date L:137 M:257 W: Feature value mis-spelled or invalid, <2215 Name/Key for SEQ ID#:3 L:175 M:257 W: Feature value mis-spelled or invalid, <2215 Name/Key for SEQ ID#:4 L:185 M:257 W: Feature value mis-spelled in invalid, <2217 Name/Key for SEQ ID#:5